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PROGRAMS FOR HAYMAN'S APPROACH - DIALLEL ANALYSIS



ICRISAT

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PROGRAMS FOR HAYMAN'S APPROACH - DIALLEL ANALYSIS

GENSTAT programs for Hayman's approach of diallel analysis in two parts have been developed. The first program will carryout the analysis of variance. In this technique, the total sum of squares is partitioned into various components, namely a (additive), b (non-additive, which is further sub-divided into b_1, b_2 and b_3), c (maternal) and d (reciprocal differences other than c).

This program will also test for homogeneity of variances in the non-additive component interactions and an indication of appropriate interaction term to be taken for testing b_1, b_2 and b_3 is given. Similarly, a test for homogeneity of variances of all other component interactions such as additive (a), non-additive (b), maternal (c) and reciprocal differences (d), will be carried out.

The second program is written to get the estimation of components of variation. This program does the testing of the significance of genotypic differences, estimation of variances and covariances, W_r-V_r graph, estimation of different components of variation such as E(expected environmental component of variation), D (variation due to additive effect), F (the mean of "Fr" over the arrays, H_1, H_2 (component of variation due to the dominance effect of genes), h_2 (dominance effect) and standard errors for all these components. Further it also calculates the

mean degree of dominance, the coefficient of correlation between $VR+VR$ (Parental order of dominance) and Y_r (Parental measurement), heritability and estimation of most dominant and recessive parent.

An example data from a 8×8 diallel where 8 parents were involved in a diallel producing 28 F_1 's and 28 reciprocals, from "Biometrical methods in Quantitative Genetic Analysis " by Singh and Chaudhary, is analysed using these programs and output is annexed.

PROGRAM HAYMAN'S DIALLEL APPROACH
(HAYMANI.PRO)

```
'REPE/NUNN=300,NID=300' HAYMAN1
'UNIT' $ 256
'SCAL' NR=4: N=8 : NR1
'CALC' NR1=NR+1
'FACT' MALE $ N=(1...N)32
'FACT' FEMY $ N=32(1...N)
'FACT' REP $ NR=8(1...NR)8
'FACT' M $ 8=(1...8)8
'FACT' F $ 8=8(1...8)
'SCAL' T,N2
'CALC' T=N*N
'CALC' N2=((N*N-1)/2)+N
'R'
'INPU' 2
'READ' YLD
'INPU' 1
'VARI' V(1...NR),TOTAL $ T
'VARI' SS(1...NR1),V1,V2 $ 8
,,
                                Total of raw data over replications
                                stored in identifier YT
,,
'FOR' K=1...NR; YT=V(1...NR)
'FOR' I=1...N: J=1...N
'REST' YLD $ REP,FEMY,MALE=K,I,J
'SCAL' IJ
'CALC' IJ=J+N*(I-1)
'CALC' ELEM(YT;IJ)=SUM(YLD)
'REPE'
'REPE'
'REPE'
'CALC' TOTAL=VSUM(V(1...NR))
'FOR' YT=V(1...NR),TOTAL; XX=SS(1...NR1)
'VARI' MT,FT,PT,Y1,Y2,Y3,Y4 $ N
'FOR' I=1...N
'REST' YT $ M=I
'CALC' ELEM(MT;I)=SUM(YT)
'REPE'
'REST' YT
'FOR' I=1...N
'REST' YT $ F=I
'CALC' ELEM(FT;I)=SUM(YT)
'REPE'
'REST' YT
'matr' YRSP,X1,X2,YRSM $ N,N
'EQUA' X1=YT
'CALC' X2=TRANS(X1) : YRSP=(X1+X2) : YRSM=(X1-X2)
'DIAG' P $ N
'CALC' P=X1
'EQUA' PT=P
'CALC' Y1=MT+FT : Y2=MT-FT : Y3=Y1-N*PT : Y4=Y1-2*PT
```

```

'SCAL' Y3S,TS,SS,SS1,SS2,SS4,SSA,SSB1,SSB,SSB2,SSB3,SSC,SSD,TSS
'CALC' TS=SUM(YT) : Y3S=SUM(Y3)
'CALC' SS=(1/(2*N))*SUM(Y1*Y1) : SS1=(2/(N*N))*SUM(TS*TS)
'CALC' SSA=SS-SS1
'CALC' SS=SUM(PT) : SS1=TS-N*(SS) : SS1=SS1*SS1
'CALC' SSB1=SS1/(N*N*(N-1))
'CALC' SS=SUM(Y3*Y3)/(N*(N-2)) : SS1=Y3S*Y3S/(N*N*(N-2))
'CALC' SSB2=SS-SS1
'CALC' SS=SUM(YRSP*YRSP)/4 : SS1=SUM(PT*PT)
'CALC' SS2=SUM(Y4*Y4)/(2*(N-2))
'CALC' SS4=TS-SUM(PT) : SS4=SS4*SS4/((N-1)*(N-2))
'CALC' SSB3=SS-SS1-SS2+SS4
'CALC' SS1=SUM(Y1*Y1)/(2*N)
'CALC' SS2=(TS*TS)/(N*N)
'CALC' SSB=SS-SS1+SS2
'CALC' SSC=(1/(2*N))*SUM(Y2*Y2)
'CALC' SS=SUM(YRSM*YRSM)/4 : SS1=SUM(Y2*Y2)/(2*N)
'CALC' SSD=SS-SS1
'CALC' TSS=SUM(YT*YT)-((TS*TS)/(N*N))
'EQUA' XX=SSA,SSB1,SSB2,SSB3,SSB,SSC,SSD,TSS
'REPE'
'CALC' SS(NR1)=SS(NR1)/NR
'SCAL' DF(1..16),S(1..16),H(1..16)
'CALC' DF(1)=N-1: DF(2)=1: DF(3)=N-1: DF(4)=N*(N-3)/2
'CALC' DF(5)=N*(N-1)/2 : DF(6)=N-1: DF(7)=(N-1)*(N-2)/2
'CALC' DF(8)=(N*N)-1 : DF(9)=(N-1)*(NR-1)
'CALC' DF(10)=1*(NR-1) : DF(11)=(N-1)*(NR-1)
'CALC' DF(12)=(N*(N-3)/2)*(NR-1) : DF(13)=(N*(N-1)/2)*(NR-1)
'CALC' DF(14)=(N-1)*(NR-1): DF(15)=((N-1)*(N-2)/2)*(NR-1)
'CALC' DF(16)=(N*N-1)*(NR-1)
'CALC' V1=VSUM(SS(1..NR))
'CALC' V2=V1-SS(NR1)
'EQUA' S(1..16)=SS(NR1),V2
'CALC' M(1)=S(1)/DF(1) : M(2)=S(2)/DF(2) : M(3)=S(3)/DF(3) : M(4)=S(4)/DF(4)
'CALC' M(5)=S(5)/DF(5) : M(6)=S(6)/DF(6) : M(7)=S(7)/DF(7)
'CALC' M(8)=S(8)/DF(8) : M(9)=S(9)/DF(9) : M(10)=S(10)/DF(10)
'CALC' M(11)=S(11)/DF(11) : M(12)=S(12)/DF(12) : M(13)=S(13)/DF(13)
: M(14)=S(14)/DF(14) : M(15)=S(15)/DF(15) : M(16)=S(16)/DF(16)
'HEAD' h(1)=' 'a ' '
: h(2)=' 'b1 ' '
: h(3)=' 'b2 ' '
: h(4)=' 'b3 ' '
: h(5)=' 'b ' '
: h(6)=' 'c ' '
: h(7)=' 'd ' '
: h(8)=' 'Total ' '
: h(9)=' 'a x blocks ' '
: h(10)=' 'b1 x blocks ' '
: h(11)=' 'b2 x blocks ' '
: h(12)=' 'b3 x blocks ' '
: h(13)=' 'b x blocks ' '
: h(14)=' 'c x blocks ' '
: h(15)=' 'd x blocks ' '
: h(16)=' 'Total x blocks ' '
'head' h=' '*** Analysis of Variance in Hayman's Analysis *** ' '

```

```

'prin' h
'LINE' 1
'head' h1='Source          d.f.          S.S.          M.S. '
'prin' h1 'line' 1
'prin/c,labr=1,labc=1' h(1),DF(1),S(1),M(1)      $ 0,24,2(12.3)
'prin/c,labr=1,labc=1' h(2),DF(2),S(2),M(2)      $ 0,23,2(12.3)
'prin/c,labr=1,labc=1' h(3),DF(3),S(3),M(3)      $ 0,23,2(12.3)
'prin/c,labr=1,labc=1' h(4),DF(4),S(4),M(4)      $ 0,23,2(12.3)
'prin/c,labr=1,labc=1' h(5),DF(5),S(5),M(5)      $ 0,24,2(12.3)
'prin/c,labr=1,labc=1' h(6),DF(6),S(6),M(6)      $ 0,24,2(12.3)
'prin/c,labr=1,labc=1' h(7),DF(7),S(7),M(7)      $ 0,24,2(12.3)
'prin/c,labr=1,labc=1' h(8),DF(8),S(8)           $ 0,20,2(12.3)
'prin/c,labr=1,labc=1' h(9),DF(9),S(9),M(9)      $ 0,15,2(12.3)
'prin/c,labr=1,labc=1' h(10),DF(10),S(10),M(10) $ 0,14,2(12.3)
'prin/c,labr=1,labc=1' h(11),DF(11),S(11),M(11) $ 0,14,2(12.3)
'prin/c,labr=1,labc=1' h(12),DF(12),S(12),M(12) $ 0,14,2(12.3)
'prin/c,labr=1,labc=1' h(13),DF(13),S(13),M(13) $ 0,15,2(12.3)
'prin/c,labr=1,labc=1' h(14),DF(14),S(14),M(14) $ 0,15,2(12.3)
'prin/c,labr=1,labc=1' h(15),DF(15),S(15),M(15) $ 0,15,2(12.3)
'prin/c,labr=1,labc=1' h(16),DF(16),S(16)      $ 0,11,2(12.3)

```

Testing for Homogeneity of blocks

```

' SCAL' NC=3 :PROBLEV=.05
' VARI' NDF,MS $ 3
' EQUA' NDF=DF(10,11,12) : MS=M(10,11,12)
' SCAL' Q,PROB,NC1
' CALC' NC1=NC-1
' CALC' Q=1+(SUM(1/NDF)-1/SUM(NDF))/3/NC1
        : Q=(SUM(NDF)*LOG(SUM(NDF*MS)/SUM(NDF))-SUM(NDF*LOG(MS)))/Q
        : PROB=1-CPROB(Q;NC1)
' JUMP' LCOMB*(PROB.GT.PROBLEV)
' CAPT'

```

Note:

Mean squares were found Heterogeneous, hence
b1,b2 and b3 are tested against
each interaction

```

' JUMP' L2
' LABE' LCOMB
' CAPT'

```

Note:

Mean squares were foundd homogeneous, hence
b1, b2 and b3 are tested against
b x blocks interaction.

```

' LABE' L2

```

Testing for Homogeneity of other
a,b,c and d interactions

```

'SCAL' NC=4
'VARI' NDF1,MS1 $ 4
'EQUA' NDF1-DF(9,13,14,15) : MS1-M(9,13,14,15)
'SCAL' Q,PROB,NC1
'CALC' NC1=NC-1
'CALC' Q=1+(SUM(1/NDF1)-1/SUM(NDF1))/3/NC1
      : Q=(SUM(NDF1)*LOG(SUM(NDF1*MS1)/SUM(NDF1))-SUM(NDF1*LOG(MS1)))/Q
      : PROB=1-CPROB(Q;NC1)
'JUMP' LCOMB1*(PROB.GT.PROBLEV)
'CAPT'
''

```

Note:

Mean squares were found Heterogeneous, hence
a,b,c and d are tested against
each of their interactions
''

```

'JUMP' L22
'LABE' LCOMB1
'CAPT'
''

```

Note:

Mean squares were found homogeneous, hence
a,b,c and d are tested against
pooled mean squares
''

```

'line' 1
'SCAL' PMS,PDF,PSS
'CALC' PDF=DF(9)+DF(13)+DF(14)+DF(15)
'CALC' PSS=S(9)+S(13)+S(14)+S(15)
'CALC' PMS=PSS/PDF
'HEAD' h(17)='' Pooled MS ''
'head' h(18)='' degrees of freedom ''
'prin/c,labr=1,labc=1' h(17),PMS $ 0,15.4
'prin/c,labr=1,labc=1' h(18),PDF $ 0,15
'LABE' L22
'RUN'
'CLOSE'
'STOP'

```

OUTPUT FROM PROGRAM 1

| IDENTIFIER | MINIMUM | MEAN | MAXIMUM | VALUES | MISSING |
|------------|---------|-------|---------|--------|---------|
| YLD | 40.64 | 90.75 | 142.84 | 256 | 0 |

*** Analysis of Variance in Hayman's Analysis ***

| Source | d.f. | S.S. | M.S. |
|----------------|------|------------|----------|
| a | 7 | 15222.500 | 2174.643 |
| b1 | 1 | 1368.045 | 1368.045 |
| b2 | 7 | 13532.469 | 1933.210 |
| b3 | 20 | 73341.000 | 3667.050 |
| b | 28 | 88241.500 | 3151.482 |
| c | 7 | 310.359 | 44.337 |
| d | 21 | 1149.456 | 54.736 |
| Total | 63 | 104924.250 | |
| a x blocks | 21 | 1678.688 | 79.938 |
| b1 x blocks | 3 | 172.459 | 57.486 |
| b2 x blocks | 21 | 2152.492 | 102.500 |
| b3 x blocks | 60 | 7088.625 | 118.144 |
| b x blocks | 84 | 9414.125 | 112.073 |
| c x blocks | 21 | 1884.185 | 89.723 |
| d x blocks | 63 | 8782.855 | 139.410 |
| Total x blocks | 189 | 21759.375 | |

Note:

Mean squares were found homogeneous, hence
b1, b2 and b3 are tested against
b x blocks interaction.

Mean squares were found homogeneous, hence
a, b, c and d are tested against
pooled mean squares

| | | |
|--------------------|----------|-----|
| Pooled MS | 115.1315 | |
| degrees of freedom | | 189 |

212 'CLOSE'

***** END OF HAYMAN1. MAXIMUM OF 53686 DATA UNITS
USED AT LINE 36 (273994 LEFT)

**PROGRAM1 FOR HAYMANS' DIALLEL APPROACH
(HAYMAN2.PRO)**

```
'REPE/NUNN=500,NID=500' DIALLEL
'SCAL' DSIZE=8 : NREP=4 : NV=1 : MF,NOBS,DSIZ3
'CALC' MF=DSIZE*DSIZE : NOBS=MF*NREP : DSIZ3=DSIZE*NREP
'INTE' NUMVAR=1...NV
'R'
'UNIT' $ NOBS
'FACT' REP $ NREP=DSIZE!(1...NREP)DSIZE
'FACT' MALE $ DSIZE=DSIZ3!(1...DSIZE)
      : FEMALE $ DSIZE=(1...DSIZE)DSIZ3
'FACT' CRS $ MF=(1...8)NREP,(9,10...16)NREP,(17,18...24)NREP,
      (25,26...32)NREP,(33,34...40)NREP,
      (41,42...48)NREP,(49,50...56)NREP,(57,58...64)NREP
'INPU' /RECL=132' 2
'READ' /P' V(NUMVAR)
'INPU' 1
'R'
'VARI' MFTOT $ MF
'FOR' YSET=V(NUMVAR)
'FOR' I=1...DSIZE : J=1...DSIZE
'REST' YSET $ MALE,FEMALE=I,J
'SCAL' IJ,JJ
'CALC' IJ=J+DSIZE*(I-1)
'CALC' JJ=SUM(YSET)
'COPY' MFTOT $IJ=JJ
'REPE' :
'REST' YSET
'BLOC' REP/CRS
'TREA' CRS
'ANOVA/PR=00010' YSET; OUT=XOUT
'EXTR' XOUT ; REP $ SS=REPSS ; DF=PDF
'EXTR' XOUT ; REP.CRS $ SS=ESS ; DF=EDF
,,
```

DIALLEL PROGRAM STARTS FROM HERE

```
,,
'SCAL' WR(1...DSIZE),PRT(1...DSIZE),WR1M,VWR,VVR,COVVV,
      CC,MDD,GEF,PDR,MFR,GGN,COR2,COR2SQ,DD,FF,H1H,H2H,DOM2,
      E2,ERROR,D,F,HH1,HH2,DOMEFF,N2,N3,
      N4,N5,SED,SEF,SEH1,SEH2,SEDOM,SEE,HERIT,VD,
      VR1,VA1,WD,T1,T2,TT,YD,YRR,
      GTOTAL,PRM,GM,TDF,EDF,RDF
'MATRI' X3,X1,X2 $ DSIZE,DSIZE
'EQUA' X1=MFTOT
'CALC' X2=TRANS(X1) : X3=(X1+X2)/(2*NREP)
'DIAGMAT' PR $ DSIZE
'CALC' PR=X1/NREP
'VARI' VM,MVA,PRT,PARENTS,P(1...DSIZE) $ DSIZE
'EQUA' P(1...DSIZE)=X3 'SYMM' MEANS $ DSIZE
'CALC' MEANS=X3 'EQUA' PARENTS=PR
,,
```

Estimation of Variances and Covariances
of parents and arrays

```

''
'SCAL' PRS 'CALC' PRS=SUM(PARENTS)
'SCAL' PRV,VR(1...DSIZE)
'CALC' PRV,VR(1...DSIZE)=VAR(PARENTS,P(1...DSIZE))
'SCAL' PRTOT(1...DSIZE) 'CALC' PRTOT(1...DSIZE)=SUM(P(1...DSIZE))
'SCAL' PRMN(1...DSIZE) 'CALC' PRMN(1...DSIZE)=MEAN(P(1...DSIZE))
'EQUA' PRT=PRTOT(1...DSIZE)
'CALC' GTOTAL=SUM(PRT) : PRM=MEAN(PARENTS)
'CALC' GM=GTOTAL/DSIZE 'EQUA' VM=PRMN(1...DSIZE) : MVA=VR(1...DSIZE)
'HEAD' H1=' *** MEAN DATA OVER REPLICATIONS AND RECIPROCALLS *** '
'HEAD' H2=' TOTAL' : H3=' MEAN ' : H4=' PARENTAL MEAN'
'HEAD' H5=' VARIANCE OF PARENTS'
'PRIN' H1
'PRIN/ LHM=5,LABC=1' MEANS $ 10.3 'LINE' 1
'PRIN/C, LABR=1,LABC=1' H2,PRTOT(1...DSIZE) $ 0,13.3,(10.3)DSIZE
'PRIN/C, LABR=1,LABC=1' H3,PRMN(1...DSIZE) $ 0,14.3,(10.3)DSIZE 'LINE' 1
'PRIN/C, LABR=1,LABC=1' H4,PRM $ 0,20.3
'PRIN/C, LABR=1,LABC=1' H5,PRV $ 0,20.3
''

```

variance of the mean arrays

```

''
'SCAL' VMA 'CALC' VMA=VAR(VM) 'SCAL' MVA1 'CALC' MVA1=MEAN(MVA)
''

```

Covariance between parents and off-spring = Vr.

```

''
'VARI' FR,VRVR1,VRVR2,VR,V1(1...DSIZE),FITTED,VRI $ DSIZE
'CALC' V1(1...DSIZE)=P(1...DSIZE)*PARENTS
'CALC' PRT(1...DSIZE)=(PRTOT(1...DSIZE)*PRS)/DSIZE
'CALC' VR(1...DSIZE)=(SUM(V1(1...DSIZE))-PRT(1...DSIZE))/(DSIZE-1)
'EQUA' VR=VR(1...DSIZE) 'CALC' : VRVR1=VR+MVA
: VRVR2=VR-MVA : VR1M=MEAN(VR)
'VARI' ARRAY $ DSIZE=1...DSIZE
'HEAD' H6=
''

```

** ARRAY VARIANCES AND COVARIANCES **

```

-----
ARRAY      VR      VR      VR-VR      VR+VR      YR
-----
'HEAD' H66=
''

```

```

-----
'PRIN' H6
'PRIN/P,LABR=1,LABC=1' ARRAY,VR,MVA,VRVR2,VRVR1,PARENTS $ 5,(10.3)5
'PRIN/C,LABR=1,LABC=1' H3,VR1M,MVA1,PRM $ 0,10.3,10.3,30.3
'PRIN' H66
''

```

Difference between the mean of parents and the mean of their
nxn progeny

```

''
'SCAL' MLO 'CALC' MLO=((GM-PRS)/DSIZE)*((GM-PRS)/DSIZE)
''

```

TESTING THE VALIDITY OF HYPOTHESIS

```

''
'CALC' VVR=VAR(VR) : VVR=VAR(MVA)
: COVVV=(SUM(VR*MVA)-(MVA1*VR1M))/(DSIZE-1)

```

VR-VR GRAPH

```

''
'HEAD' HX=''' VR '' : HY='''VR'' : HH='''SP''
''

```

CALCULATION OF INTERCEPT

```

''
'HEAD' H7=''' INTERCEPT VALUE A= ''
'SCAL' B,INTCPT
'TERMS' VR,MVA
'Y' VR
'FIT' MVA; COEF=CC1; PVAL=FITTED
'COPY' INTCPT = CC1 $ 1 : B =CC1 $ 2
'GRAPH/ATX=HX,ATY=HY,NRF=30,NCF=40' FITTED,VR;MVA $ HH
'CAPT'
''

```

*** ESTIMATION OF COMPONENTS OF VARIATION ***

```

''
'HEAD' H8=''' EXPECTED ENVIRONMENTAL COMPONENT OF VARIATION (E) = ''
'HEAD' H9=''' VARIATION DUE TO ADDITIVE EFFECT (D) = ''
'HEAD' H10=''' MEAN OF 'FR' OVER ARRAYS (F) = ''
'HEAD' H11=''' COMPONENT OF VARIATION DUE TO DOMINANCE EFF. OF GENES =''
'HEAD' H12=''' DOMINANCE EFFECT (h2) = ''
'CALC' ERROR=((ESS+REPSS)/(EDF+RDF))/NREP
'CALC' D=PRV-ERROR : F=(2*PRV)-(4*VR1H)-(2*(DSIZE-2)*ERROR/DSIZE)
'CALC' HH1=PRV-(4*VR1H)+(4*MVA1)-((3*DSIZE-2)*ERROR/DSIZE)
'CALC' HH2=(4*MVA1)-(4*MVA)-(2*ERROR)
'CALC' DOMEFF=(4*MLO)-(4*DSIZE-1)*ERROR/(DSIZE*DSIZE)
'PRIN/C,LABR=1,LABC=1' H8,ERROR $ 0,20.3 : H9,D $ 0,20.3 : H10,F $ 0,20.3
'PRIN/C' H11,HH1 $ 6.3 : H11,HH2 $ 6.3
'PRIN/C,LABC=1,LABR=1' H12,DOMEFF $ 0,8.3 'LINE' 1
''

```

CALCULATION OF STANDARD ERRORS FOR TESTING COMPONENTS OF VARIATION

```

''
'SCAL' COMLTPL 'CALC' COMLTPL=(VAR(WRVR2))/2
''

```

CALCULATION OF SPECIFIC MULTIPLIERS

```

''
'CALC' N2=DSIZE*DSIZE: N3=DSIZE*DSIZE*DSIZE: N4=N2*N2 : N5=N4*DSIZE
'CALC' DD=(N5+N4)/N5 : FF=((4*N5)+(20*N4)-(16*N3)+(16*N2))/N5
'CALC' H1H=(N5+(41*N4)-(12*N3)+(4*N2))/N5 : E2=N4/N5
'CALC' H2H=(36*N4)/N5 : DOM2=((16*N4)+(16*N2)-(32*DSIZE)+16)/N5
'CALC' SED=SQRT(DD*COMLTPL) : SEF=SQRT(FF*COMLTPL)
'CALC' SEH1=SQRT(H1H*COMLTPL) : SEH2=SQRT(H2H*COMLTPL)
'CALC' SEDOM=SQRT(DOM2*COMLTPL) : SEE=SQRT(E2*COMLTPL)
'HEAD' HEAD2=''' SE(D) = '' : HEAD3=''' SE(F) = '' : HEAD4=''' SE(H1) =''
'HEAD' HEAD5=''' SE(H2) = '' : HEAD6=''' SE(DOMEFF) = ''
'HEAD' HEAD7=''' SE(E) = '' 'LINE' 1
'CAPT'
''*** STANDARD ERRORS OF DIFFERENT PARAMETERS *** '' 'LINE' 1
'PRIN/C,LABR=1,LABC=1' HEAD2,SED $ 0,10.3 : HEAD3,SEF $ 0,10.3
'PRIN/C,LABR=1,LABC=1' HEAD4,SEH1 $ 0,10.3: HEAD5,SEH2 $ 0,10.3
'PRIN/C,LABR=1,LABC=1' HEAD6,SEDOM $ 0,10.3 : HEAD7,SEE $ 0,10.3 'LINE' 1
'HEAD' HZ1='''MEAN DEGREE OF DOMINANCE = ''

```

```

'HEAD' HZ2='PROPORTION OF GENES WITH + AND - EFFECTS IN THE PARENTS = ''
'HEAD' HZ3='PROPORTION OF DOMINANT AND RECESSIVE GENES IN PARENTS = ''
'HEAD' HZ4=
'COEFFICIENT OF CORRELATION (SMALL R) BETWEEN
THE PARENTAL ORDER OF DOMINANCE (VR+VR) AND
PARENTAL MEASUREMENT YR = ''
'HEAD' HZ5='PREDICTION FOR MEASUREMENT OF DOMINANT AND RECESSIVE PARENTS = ''
'HEAD' HZ6='NO. OF GROUPS OF GENES WHICH CONTROL THE CHARACTER AND
EXHIBIT THE DOMINANCE = ''
'HEAD' HZ7='THE COVARIANCE OF ADDITIVE DOMINANCE EFFECTS IN A SINGLE ARRAY''
'HEAD' HZ8='MEAN OF FR '' ; HZ9='HERITABILITY ''
'CALC' MDD=SQRT(HH1/D) : GEF=HH2/(4*HH1)
'CALC' PDR=(SQRT(4*D*HH1)+F)/(SQRT(4*D*HH1)-F)
'SYMM' COR1 $ 2 'DSSP' COR $ VRVR1,PARENTS
'SSP/PRINT=2' COR
'CALC' COR1=CORMAT(COR) 'EQUA' COR2=COR1 $ X,1,2X
'CALC' COR2SQ=COR2*COR2
'CALC' FR=2*(PRV-VR1M*MVA1-VRVR1)-2*(DSIZE-2)*(ERROR/DSIZE)
'CALC' MFR=MEAN(FR) : GGN=DOMEFF/HH2 'LINE' 1
'PRIN/C,LABR=1,LABC=1' HZ1,MDD $ 0,5.3 : HZ2,GEF $ 0,5.3
'PRIN/C,LABR=1,LABC=1' HZ3,PDR $ 0,5.3 'LINE' 1
'PRIN/C,LABR=1,LABC=1' HZ4,COR2 $ 0,5.3 'LINE' 1
'PRIN/C,LABR=1,LABC=1' HZ5,COR2SQ $ 0,5.3 : HZ6,GGN $ 0,5.3 : HZ7
'PRIN/C,LABR=1,LABC=1' FR $ 10.3 : HZ8,MFR $ 0,8.3 'LINE' 1

```

CALCULATION OF HERITABILITY

```

''
'SCAL' HERIT1
'CALC' HERIT=(D/2)+(HH1/2)-(HH2/2)-(MFR/2)
'CALC' HERIT1=(D/2)+(HH1/2)-(HH2/4)-(MFR/2)+ERROR
'CALC' HERIT=HERIT/HERIT1
'PRIN/C,LABR=1,LABC=1' HZ9,HERIT $ 0,8.3
'CAPT'
''
** ESTIMATION OF MOST DOMINANT AND RECESSIVE PARENT **
''
'CALC' TT=(PRV+SQRT((PRV*PRV)-(4*PRV*(VR1M-MVA1))))/(2*PRV)
'CALC' T1=TT-1 : T2=1-T1
'CALC' VD=PRV*(T2*T2) : WD=PRV*T2 : VR1=PRV*(T1*T1)
'CALC' WA1=PRV*T1 : YD=PRM+B*((WD+VD)-(VR1M*MVA1))
'CALC' YRR=PRM+B*((WA1+VR1)-(VR1M*MVA1))
'HEAD' HL1='VALUE OF COMPLETELY DOMINANT PARENT YD = ''
'HEAD' HL2='VALUE OF COMPLETELY RECESSIVE PARENT YR = ''
'PRIN/C,LABR=1,LABC=1' HL1,YD $ 0,10.3 : HL2,YRR $ 0,10.3
'REPE'
'RUN'
'CLOSE'
'STOP'

```

OUTPUT FROM PROGRAM2

| | | | | | |
|------------|---------|-------|---------|--------|---------|
| IDENTIFIER | MINIMUM | MEAN | MAXIMUM | VALUES | MISSING |
| V(1) | 40.64 | 90.75 | 142.84 | 256 | 0 |

***** ANALYSIS OF VARIANCE *****

VARIATE: V(1)

| SOURCE OF VARIATION | DF | SS | SSX | MS | VR |
|------------------------------|-------|----------|--------|--------|--------|
| REP STRATUM | 3 | 1037.0 | 0.81 | 345.7 | |
| REP.CRS STRATUM | | | | | |
| CRS | 63 | 104924.2 | 82.15 | 1665.5 | 14.467 |
| RESIDUAL | 189 | 21758.5 | 17.04 | 115.1 | |
| TOTAL | 252 | 126682.6 | 99.19 | 502.7 | |
| GRAND TOTAL | 255 | 127719.7 | 100.00 | | |
| GRAND MEAN | 90.75 | | | | |
| TOTAL NUMBER OF OBSERVATIONS | 256 | | | | |

*** MEAN DATA OVER REPLICATIONS AND RECIPROCAL ***

| | | | | | | | | | |
|---------------------|---------|---------|---------|---------|---------|---------|---------|---------|--|
| 1 | 85.645 | | | | | | | | |
| 2 | 83.850 | 98.260 | | | | | | | |
| 3 | 96.368 | 108.065 | 74.070 | | | | | | |
| 4 | 117.030 | 88.740 | 101.660 | 91.640 | | | | | |
| 5 | 115.790 | 101.410 | 91.275 | 84.338 | 54.100 | | | | |
| 6 | 68.693 | 72.280 | 100.230 | 107.230 | 86.058 | 100.390 | | | |
| 7 | 108.453 | 92.730 | 109.993 | 61.050 | 80.343 | 118.640 | 90.960 | | |
| 8 | 50.680 | 84.398 | 116.568 | 46.798 | 94.070 | 56.990 | 121.720 | 82.000 | |
| TOTAL | 726.508 | 729.733 | 798.227 | 698.485 | 707.383 | 710.510 | 783.888 | 653.223 | |
| MEAN | 90.813 | 91.217 | 99.778 | 87.311 | 88.423 | 88.814 | 97.986 | 81.653 | |
| PARENTAL MEAN | 84.633 | | | | | | | | |
| VARIANCE OF PARENTS | 224.986 | | | | | | | | |

**** ARRAY VARIANCES AND COVARIANCES ****

| ARRAY | VR | VR | VR-VR | VR+VR | YR |
|-------|----------|---------|----------|---------|---------|
| 1 | -124.142 | 546.907 | -671.049 | 422.766 | 85.645 |
| 2 | -98.412 | 129.310 | -227.722 | 30.897 | 98.260 |
| 3 | 97.340 | 171.925 | -74.585 | 269.266 | 74.070 |
| 4 | 39.070 | 549.498 | -510.428 | 588.568 | 91.640 |
| 5 | 155.804 | 317.612 | -161.809 | 473.416 | 54.100 |
| 6 | 43.122 | 455.954 | -412.833 | 499.076 | 100.390 |
| 7 | 44.363 | 428.021 | -383.658 | 472.384 | 90.960 |
| 8 | -160.305 | 824.103 | -984.408 | 663.797 | 82.000 |
| MEAN | -0.395 | 427.916 | | | 84.633 |

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******* REGRESSION ANALYSIS *******

Y-VARIATE: VR

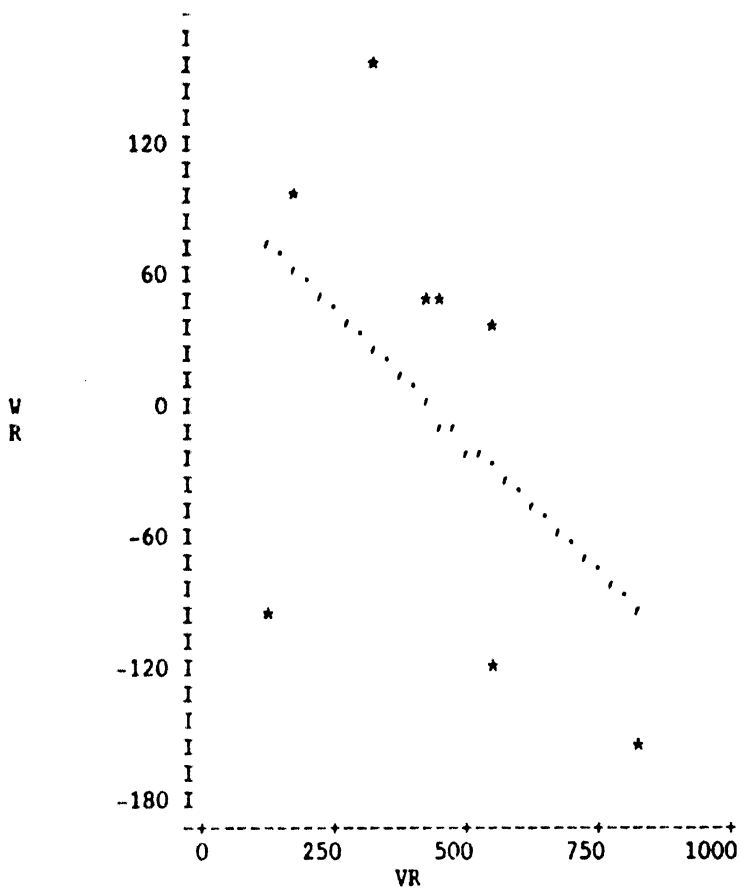
***** REGRESSION COEFFICIENTS *****

| | ESTIMATE | S.E. | T |
|----------|----------|-------|-------|
| CONSTANT | 102.5 | 86.3 | 1.19 |
| MVA | -0.241 | 0.181 | -1.33 |

***** ANALYSIS OF VARIANCE *****

| | DF | SS | MS |
|----------|----|--------|-------|
| REGRESSN | 1 | 20453 | 20453 |
| RESIDUAL | 6 | 69444 | 11574 |
| TOTAL | 7 | 89897 | 12842 |
| CHANGE | -1 | -20453 | 20453 |

PERCENTAGE VARIANCE ACCOUNTED FOR 9.9



*** ESTIMATION OF COMPONENTS OF VARIATION ***

EXPECTED ENVIRONMENTAL COMPONENT OF VARIATION (E) = 29.682
 VARIATION DUE TO ADDITIVE EFFECT (D) = 195.305
 MEAN OF 'FR' OVER ARRAYS (F) = 407.031
 COMPONENT OF VARIATION DUE TO DOMINANCE EFF. OF GENES = HH1 1856.608
 COMPONENT OF VARIATION DUE TO DOMINANCE EFF. OF GENES = HH2 1516.390
 DOMINANCE EFFECT (h2) = 135.253

*** STANDARD ERRORS OF DIFFERENT PARAMETERS ***

SE(D) = 222.038
 SE(F) = 524.656
 SE(H1) = 510.433
 SE(H2) = 444.077
 SE(DOMEFF) = 297.817
 SE(E) = 74.013

MEAN DEGREE OF DOMINANCE = 3.083
 PROPORTION OF GENES WITH + AND - EFFECTS IN THE PARENTS = 0.204
 PROPORTION OF DOMINANT AND RECESSIVE GENES IN PARENTS = 2.021

COEFFICIENT OF CORRELATION (SMALL R) BETWEEN
 THE PARENTAL ORDER OF DOMINANCE (VR+VR) AND
 PARENTAL MEASUREMENT YR = -0.155

PREDICTION FOR MEASUREMENT OF DOMINANT AND RECESSIVE PARENTS = 0.024
 NO. OF GROUPS OF GENES WHICH CONTROL THE CHARACTER AND
 EXHIBIT THE DOMINANCE = 0.089

THE COVARIANCE OF ADDITIVE DOMINANCE EFFECTS IN A SINGLE ARRAY

416.542
 1200.279
 723.542
 84.938
 315.241
 263.921
 317.307
 -65.521

MEAN OF FR 407.031

HERITABILITY 0.136

** ESTIMATION OF MOST DOMINANT AND RECESSIVE PARENT **

VALUE OF COMPLETELY DOMINANT PARENT YD = 185.647
 VALUE OF COMPLETELY RECESSIVE PARENT YR = 84.443